

Pei-kun Wang

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

260
citations

840776

11
h-index

996975

15
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19
all docs

19
docs citations

19
times ranked

129
citing authors

#	ARTICLE	IF	CITATIONS
1	The emergence of the infection of subgroup J avian leucosis virus escalated the tumour incidence in commercial Yellow chickens in Southern China in recent years. <i>Transboundary and Emerging Diseases</i> , 2019, 66, 312-316.	3.0	38
2	Diversity and evolution analysis of glycoprotein GP85 from avian leukosis virus subgroup J isolates from chickens of different genetic backgrounds during 1989-2016: Coexistence of five extremely different clusters. <i>Archives of Virology</i> , 2018, 163, 377-389.	2.1	22
3	Full-length genome sequence analysis of an avian leukosis virus subgroup J (ALV-J) as contaminant in live poultry vaccine: The commercial live vaccines might be a potential route for ALV-J transmission. <i>Transboundary and Emerging Diseases</i> , 2018, 65, 1103-1106.	3.0	20
4	Full-length genome sequence analysis of four subgroup J avian leukosis virus strains isolated from chickens with clinical hemangioma. <i>Virus Genes</i> , 2017, 53, 868-875.	1.6	19
5	Vertical transmission of ALV from ALV-J positive parents caused severe immunosuppression and significantly reduced Marek's disease vaccine efficacy in three-yellow chickens. <i>Veterinary Microbiology</i> , 2020, 244, 108683.	1.9	19
6	Analysis of the evolution and transmission dynamics of the field MDV in China during the years 1995-2020, indicating the emergence of a unique cluster with the molecular characteristics of vv+MDV that has become endemic in southern China. <i>Transboundary and Emerging Diseases</i> , 2021, 68, 3574-3587.	3.0	19
7	The Emergence of a vv + MDV Can Break through the Protections Provided by the Current Vaccines. <i>Viruses</i> , 2020, 12, 1048.	3.3	17
8	Complete genome sequencing and characterization revealed a recombinant subgroup B isolate of avian leukosis virus with a subgroup J-like U3 region. <i>Virus Genes</i> , 2017, 53, 927-930.	1.6	15
9	A novel recombinant avian leukosis virus isolated from gamecocks induced pathogenicity in Three-Yellow chickens: a potential infection source of avian leukosis virus to the commercial chickens. <i>Poultry Science</i> , 2019, 98, 6497-6504.	3.4	15
10	Full-length cDNA sequence analysis of 85 avian leukosis virus subgroup J strains isolated from chickens in China during the years 1988-2018: coexistence of 2 extremely different clusters that are highly dependent upon either the host genetic background or the geographic location. <i>Poultry Science</i> , 2020, 99, 3469-3480.	3.4	15
11	Genetic diversity of avian leukosis virus subgroup J (ALV-J): toward a unified phylogenetic classification and nomenclature system. <i>Virus Evolution</i> , 2021, 7, veab037.	4.9	15
12	Rapid detection of the common avian leukosis virus subgroups by real-time loop-mediated isothermal amplification. <i>Virology Journal</i> , 2015, 12, 195.	3.4	13
13	An outbreak in three-yellow chickens with clinical tumors of high mortality caused by the coinfection of reticuloendotheliosis virus and Marek's disease virus: a speculated reticuloendotheliosis virus contamination plays an important role in the case. <i>Poultry Science</i> , 2021, 100, 19-25.	3.4	10
14	ALV-J-contaminated commercial live vaccines induced pathogenicity in Three-Yellow chickens: one of the transmission routes of ALV-J to commercial chickens. <i>Poultry Science</i> , 2021, 100, 101027.	3.4	8
15	Recombinant subgroup B avian leukosis virus combined with the subgroup J env gene significantly increases its pathogenicity. <i>Veterinary Microbiology</i> , 2020, 250, 108862.	1.9	6
16	Reemergence of reticuloendotheliosis virus and Marek's disease virus co-infection in Yellow-Chickens in Southern China. <i>Poultry Science</i> , 2021, 100, 101099.	3.4	5
17	Two novel recombinant avian leukosis virus isolates from Luxi gamecock chickens. <i>Archives of Virology</i> , 2020, 165, 2877-2881.	2.1	4